

E. Willis

Re-run

STIC copy #11

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/522,342

DATE: 10/15/2001

TIME: 15:49:39

Input Set : A:\P1219P1-US Sequence Listing.txt

Output Set: N:\CRF3\10152001\I522342.raw

3 <110> APPLICANT: Stewart, Timothy A.  
 4 Tomlinson, Elizabeth  
 5 Goddard, Audrey  
 6 Gurney, Austin L.  
 8 <120> TITLE OF INVENTION: FIBROBLAST GROWTH FACTOR-19 (FGF-19) NUCLEIC ACIDS AND  
 9 POLYPEPTIDES AND METHODS FOR THE TREATMENT OF OBESITY  
 11 <130> FILE REFERENCE: P1219P1-US  
 13 <140> CURRENT APPLICATION NUMBER: US 09/522,342  
 14 <141> CURRENT FILING DATE: 2000-03-09  
 16 <150> PRIOR APPLICATION NUMBER: US 60/066,840  
 17 <151> PRIOR FILING DATE: 1997-11-25  
 19 <150> PRIOR APPLICATION NUMBER: US 09/158,342  
 20 <151> PRIOR FILING DATE: 1998-09-21  
 22 <150> PRIOR APPLICATION NUMBER: US 09/284,663  
 23 <151> PRIOR FILING DATE: 1999-04-15  
 25 <150> PRIOR APPLICATION NUMBER: PCT/US98/25190  
 26 <151> PRIOR FILING DATE: 1998-11-25  
 28 <160> NUMBER OF SEQ ID NOS: 5  
 30 <210> SEQ ID NO: 1  
 31 <211> LENGTH: 2137  
 32 <212> TYPE: DNA  
 33 <213> ORGANISM: Homo Sapien  
 35 <400> SEQUENCE: 1

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38	gaaacccggc	cgctaagcga	ggcctcctcc	tcccgagat	ccgaacggcc	100
40	tgggcggggg	caccccggt	gggacaagaa	gccgcgcct	gcctgcccgg	150
42	gcccggggag	ggggtgggg	ctggggccgg	aggcggggtg	tgagtgggtg	200
44	tgtgcggggg	gcggaggtt	gatgcaatcc	cgataagaaa	tgctcgggtg	250
46	tcttgggcac	ctaccctgg	ggcccgtaa	gcgctactat	ataaggctgc	300
48	cgcccgggag	ccgcgcgcgc	gtcagagcag	gagcgtgcg	tccaggatct	350
50	agggccacga	ccatcccaac	ccggcactca	cagccccgca	gcgcaccccg	400
52	gtcgcgcgcc	agcctcccg	acccccatcg	ccggagctgc	gccgagagcc	450
54	ccagggaggt	gccatgcgga	gcgggtgtgt	ggtggtccac	gtatggatcc	500
56	tggccggcct	ctggctggcc	gtggccgggc	gccccctcgc	cttctcggac	550
58	gcggggcccc	acgtgcacta	cggtcggggc	gacccccatcc	gcctgcggca	600
60	cctgtacacc	tccggccccc	acgggctctc	cagctgcttc	ctgcgcacatcc	650
62	gtgcgcagcg	cgctcgtggac	tgcgcgcggg	gccagagcgc	gcacagtttg	700
64	ctggagatca	aggcagtcgc	tctgcggacc	gtggccatca	agggcgtgca	750
66	cagcgtgcgg	tacctctgca	tgggcgccga	cggcaagatg	caggggctgc	800
68	ttcagtactc	ggaggaagac	tgtgctttcg	aggaggagat	ccgcccagat	850
70	ggctacaatg	tgtaccgatc	cgagaagcac	cgctcccg	tctccctgag	900
72	cagtgcacaaa	cagcggcagc	tgtacaagaa	cagaggcttt	cttccactct	950
74	ctcatttctc	gcccattgctg	cccatggtcc	cagaggagcc	tgaggacctc	1000
76	aggggccact	tggaaatctga	catgttctct	tgcgccctgg	agaccgacag	1050
78	catggaccca	tttgggcttg	tcaccggact	ggaggccgtg	aggagtccca	1100
80	gctttgagaa	gtaactgaga	ccatgcccgg	gcctcttcac	tgctgccagg	1150
82	ggctgtggta	cctgcagcgt	gggggacgtg	cttctacaag	aacagtccctg	1200

ENTERED

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84 agtccacggt ctgttttagct ttaggaagaa acatctagaa gttgtacata 1250
86 ttcagagttt tccattggca gtgccagttt ctagccaata gacttgtctg 1300
88 atcataacat tgtaagcctg tagcttgccc agctgctgcc tgggccccca 1350
90 ttctgctccc tcgaggttgc tggacaagct gctgcaactgt ctcagttctg 1400
92 cttgaatacc tccatcgatg gggaaactcac ttcctttgga aaaattctta 1450
94 tgtcaagctg aaattctcta attttttctc atcacttccc caggagcagc 1500
96 cagaagacag gcagtagttt taatttcagg aacaggatgat ccactctgta 1550
98 aaacagcagg taaatttcac tcaaccccat gtgggaattg atctatatct 1600
100 ctacttccag ggaccatttg cccttcccaa atccctccag gccagaactg 1650
102 actggagcag gcatggccca ccaggcttca ggagtagggg aagcctggag 1700
104 cccactcca gccctgggac aacttgagaa ttccccctga ggccagttct 1750
106 gtcattgatg ctgtcctgag aataacttgc tgtcccggtg tcacctgctt 1800
108 ccactctcca gccaccagc cctctgccca cctcacatgc ctccccatgg 1850
110 attggggcct cccaggcccc ccaccttatg tcaacctgca cttcttggtc 1900
112 aaaaatcagg aaaagaaaag atttgaagac cccaagtctt gtcaataact 1950
114 tgctgtgtgg aagcagcggg ggaagaccta gaaccctttc cccagcactt 2000
116 ggttttccaa catgatattt atgagtaatt tattttgata tgtacatctc 2050
118 ttattttctt acattattta tgcccccaaa ttatatttat gtatgtaagt 2100
120 gaggtttggt ttgtatatta aaatggagtt tgtttgt 2137
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123 <211> LENGTH: 216
124 <212> TYPE: PRT
125 <213> ORGANISM: Homo Sapien
127 <400> SEQUENCE: 2
128 Met Arg Ser Gly Cys Val Val Val His Val Trp Ile Leu Ala Gly
129 1 5 10 15
131 Leu Trp Leu Ala Val Ala Gly Arg Pro Leu Ala Phe Ser Asp Ala
132 20 25 30
134 Gly Pro His Val His Tyr Gly Trp Gly Asp Pro Ile Arg Leu Arg
135 35 40 45
137 His Leu Tyr Thr Ser Gly Pro His Gly Leu Ser Ser Cys Phe Leu
138 50 55 60
140 Arg Ile Arg Ala Asp Gly Val Val Asp Cys Ala Arg Gly Gln Ser
141 65 70 75
143 Ala His Ser Leu Leu Glu Ile Lys Ala Val Ala Leu Arg Thr Val
144 80 85 90
146 Ala Ile Lys Gly Val His Ser Val Arg Tyr Leu Cys Met Gly Ala
147 95 100 105
149 Asp Gly Lys Met Gln Gly Leu Leu Gln Tyr Ser Glu Glu Asp Cys
150 110 115 120
152 Ala Phe Glu Glu Glu Ile Arg Pro Asp Gly Tyr Asn Val Tyr Arg
153 125 130 135
155 Ser Glu Lys His Arg Leu Pro Val Ser Leu Ser Ser Ala Lys Gln
156 140 145 150
158 Arg Gln Leu Tyr Lys Asn Arg Gly Phe Leu Pro Leu Ser His Phe
159 155 160 165
161 Leu Pro Met Leu Pro Met Val Pro Glu Glu Pro Glu Asp Leu Arg
162 170 175 180
164 Gly His Leu Glu Ser Asp Met Phe Ser Ser Pro Leu Glu Thr Asp

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```
165                               185                               190                               195
167 Ser Met Asp Pro Phe Gly Leu Val Thr Gly Leu Glu Ala Val Arg
168                               200                               205                               210
170 Ser Pro Ser Phe Glu Lys
171                               215
173 <210> SEQ ID NO: 3
174 <211> LENGTH: 26
175 <212> TYPE: DNA
176 <213> ORGANISM: Artificial Sequence
178 <220> FEATURE:
179 <223> OTHER INFORMATION: Synthetic oligonucleotide probe
181 <400> SEQUENCE: 3
182 atccgcccag atggctacaa tgtgta 26
184 <210> SEQ ID NO: 4
185 <211> LENGTH: 22
186 <212> TYPE: DNA
187 <213> ORGANISM: Artificial Sequence
189 <220> FEATURE:
190 <223> OTHER INFORMATION: Synthetic oligonucleotide probe
192 <400> SEQUENCE: 4
193 ccagtccggt gacaagccca aa 22
195 <210> SEQ ID NO: 5
196 <211> LENGTH: 42
197 <212> TYPE: DNA
198 <213> ORGANISM: Artificial Sequence
200 <220> FEATURE:
201 <223> OTHER INFORMATION: Synthetic oligonucleotide probe
203 <400> SEQUENCE: 5
204 gcctcccgt ctcctgagc agtgccaaac agcggcagtg ta 42
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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/522,342

DATE: 10/15/2001

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Input Set : A:\P1219P1-US Sequence Listing.txt

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